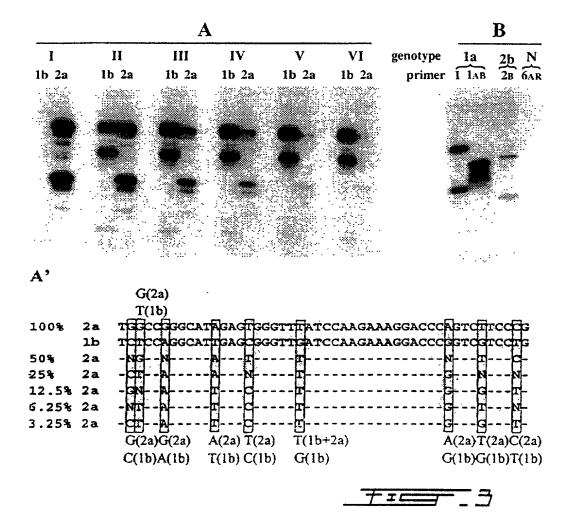
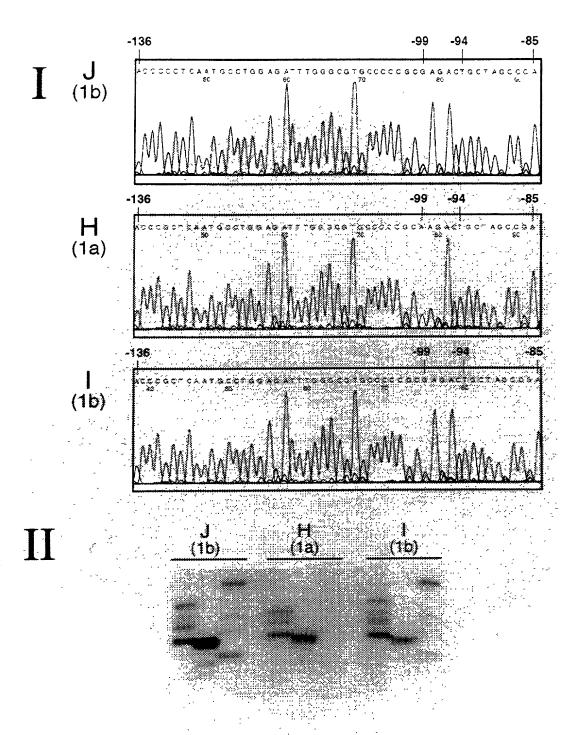


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1AB 1 1BR 1AB 1 1BR 1AB 1 1BR

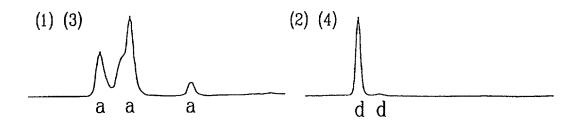
Expected base extension

					5/	10				
ATC		ATG			Acu	၁သ၁ သာ	Z L	F # 57	7 7 7 7	
m	\vdash	0	2	r(SEQ ID NO:16)		0		Ŋ	2	C GTCCACATCCT(SEQ ID NO:17)
Primer →T CAXX	Primer →T X-XX	Primer →X-X	Primer →T CXX	A GTCCACATCCT		Primer →X X	Primer →G XX	Primer → G CAGGX-X	Primer →G CXX	GICCACATCC
ATC	ATG	ACG	TCG	WT		ATC	ATG	ACG	TCG	MU

2

2 2 2	<pre>xtcatactaxtccatgagacattttcg 5' L63 + PRIMER (1) (SEQ ID NO:18) AGACAGTATGATCAGGTACTTTAGACACCTGTATTTCG 5' L63 + PRIMER (2) (SEQ ID NO:20) x GAGTATCTTTAGACACCTGTATTTCG 5' L63P + PRIMER (2) (SEQ ID NO:20) AGACAGTATGATCAGGTACCTGTGTGATAAAGC 3' PRO63L(G)-R (SEQ ID NO:21)</pre>	20)
3,	XTCATACTAXTCCATGGGTATCTTTAGACACCTGTATTTCG 5' L63P + PRIMER (3) (SEQ ID NO:22)	22)0
3,	X GGGTATCTTTAGACACCTGTATTTCG 5' L63 + PRIMER (4) (SEQ ID NO:23) AGACAGTATGATCAGGTACTCATAGAAATCTGTGGACATAAAGC 3' PRO63P(G)-R (SEQ ID NO:19)	23)

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PRIMER-SPECIFIC AND MISPAIR EXTENSION ASSAY FOR IDENTIFYING GENE VARIATION Yu-Wen Hu Serial No. 09/782,361 Filed February 13, 2001

CTGTCAACATAATTGGAAGAAATCTGAT**GACGCAGA**XX 3'PRO90M-F (3) (SEQ ID NO:28) CGACAGTTGTATTAACCTTCTTTAGACTACTGCGTCTAA---5'L90M (SEQ ID NO:29) NO:24) (2) (SEQ ID NO:26) (SEQ ID NO:30) (SEQ ID NO:25) ID NO:27) (SEQ ID L90M (SEQ 190 XXAGACTTCTGAGTCTAACCAACGTGAAATTTA 5' PRO90L-R PRO90M-F TICTGAGICTAACCAACGIGAAAITTA 5' PRO90L-R --AATCTGAAGACTCAGATTGGTTGCACTTTAAAT---3' -AATCTGATGACTCAGATTGGTTGCACTTTAAAT--3' CTGTCAACATAATTGGAAGAAATCTGAT 2 2 'n

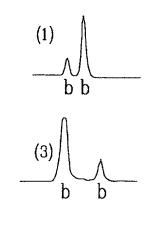
TIET 7A

NO:31)

(SEQ ID

--GACAGTTGTATTAACCTTCTTTAGACAACTGAGTCTAA--5'L90

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